

AUG 1 4 2002

TECH CENTER 1600/2900

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1644

RAW SEQUENCE LISTING PATENT APPLICATION: US/10/086,217 DATE: 07/09/2002 TIME: 16:15:10

Input Set : A:\A061cip2.app

Output Set: N:\CRF3\07092002\J086217.raw

ENTERED

3 <110> APPLICANT: MUNDY, GREGORY R. YONEDA, TOSHIYUKI <120> TITLE OF INVENTION: METHODS OF TREATING MULTIPLE MYELOMA AND MYELOMA-INDUCED BONE RESORPTION USING INTEGRIN 7 8 ANTAGONISTS 10 <130> FILE REFERENCE: A061CIP2 12 <140> CURRENT APPLICATION NUMBER: 10/086,217 C--> 13 <141> CURRENT FILING DATE: 2002-06-21 15 <150> PRIOR APPLICATION NUMBER: 09/943,659 16 <151> PRIOR FILING DATE: 2001-08-31 18 <150> PRIOR APPLICATION NUMBER: 09/805,840 19 <151> PRIOR FILING DATE: 2001-03-13 21 <150> PRIOR APPLICATION NUMBER: PCT/US99/21170 22 <151> PRIOR FILING DATE: 1999-09-13 24 <150> PRIOR APPLICATION NUMBER: 60/100,182 25 <151> PRIOR FILING DATE: 1998-09-14 27 <160> NUMBER OF SEQ ID NOS: 8 29 <170> SOFTWARE: PatentIn Ver. 2.1 31 <210> SEQ ID NO: 1 32 <211> LENGTH: 22 33 <212> TYPE: DNA 34 <213> ORGANISM: Artificial Sequence 36 <220> FEATURE: 37 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer 39 <400> SEQUENCE: 1 22 40 gctgcgcgtc accattgttc tc 43 <210> SEO ID NO: 2 44 <211> LENGTH: 22 45 <212> TYPE: DNA 46 <213> ORGANISM: Artificial Sequence 48 <220> FEATURE: 49 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer 51 <400> SEQUENCE: 2 22 52 accaccctct tgaagccttg tg 55 <210> SEQ ID NO: 3

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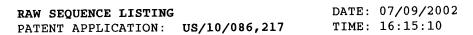
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Input Set : A:\A061cip2.app

Output Set: N:\CRF3\07092002\J086217.raw

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/086,217

DATE: 07/09/2002 TIME: 16:15:10

Input Set : A:\A061cip2.app
Output Set: N:\CRF3\07092002\J086217.raw

W--> 136 Arg Cys Asp Xaa Cys

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/086,217 DATE: 07/09/2002 TIME: 16:15:11

Input Set : A:\A06lcip2.app

Output Set: N:\CRF3\07092002\J086217.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:8; Xaa Pos. 4